

Sequence Listing

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<120> METHOD FOR MAKING MONOCLONAL ANTIBODIES AND  
CROSS-REACTIVE ANTIBODIES OBTAINABLE BY THE METHOD

<130> P1468R1 (REVISED)

<140> US 09/329,633  
<141> 1999-06-10

<150> US 60/089,253  
<151> 1998-06-12

<160> 2

<210> 1  
<211> 1799  
<212> DNA  
<213> human

<400> 1  
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gcgcacacaa aatacaccga cgatgccccga tctactttaa gggctgaaac 100  
ccacgggcct gagagactat aagagcgttc cctaccgcca tggacaacacg 150  
gggacagaac gccccggccg cttcgaaaaagg cacggcccaag 200  
gaccacggga ggcgcgggga gccaggcctg ggctccgggt ccccaagacc 250  
cttgtgctcg ttgtcgccgc ggtcctgctg ttggtctcag ctgagtctgc 300  
tctgatcacc caacaagacc tagctccccca gcagagagcg gccccacaac 350  
aaaagaggtc cagccccctca gagggattgt gtccacctgg acaccatatac 400  
tcagaagacg gtagagattg catctcctgc aaatatggac aggactatag 450  
cactcactgg aatgacacctt ttttctgttt gcgcgtgcacc aggtgtgatt 500  
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tcaacatgtt gtccccggg gagtcagagc atctgctgga accggcagaa 1000  
gctgaaaggt ctcagaggag gaggctgctg gttccagcaa atgaaggtga 1050  
tcccactgag actctgagac agtgcttcga tgactttgca gacttggtgc 1100  
cctttgactc ctgggagccg ctcatgagga agttggccct catggacaat 1150  
gagataaagg tggctaaagc tgaggcagcg ggccacaggg acaccttgta 1200  
cacgatgctg ataaagtggg tcaacaaaac cggcgagat gcctctgtcc 1250  
acaccctgct gnatgccttg gagacgctgg gagagagact tgccaagcag 1300  
aagattgagg accacttggt gagctctgga aagttcatgt atctagaagg 1350  
taatgcagac tctgccwtgt cctaagtgtg attctttca ggaagtgaga 1400  
ccttcctgg tttaccttt ttctggaaaa agcccaactg gactccagtc 1450  
agtaggaaag tgccacaatt gtcacatgac cggtaactgga agaaaactctc 1500  
ccatccaaca tcacccagtg gatgaaacat cctgtaactt ttcactgcac 1550  
ttggcattat ttttataagc tgaatgtgat aataaggaca ctatggaaat 1600  
gtctggatca ttccgtttgt gcgtactttg agatttggtt tgggatgtca 1650  
ttgttttac agcactttt tattctaatg taaatgctt atttatttat 1700  
ttgggctaca ttgtaagatc catctacaaa aaaaaaaaaa aaaaaaaaaag 1750  
ggcggccgcg actctagagt cgacctgcag aagttggcc gccatggcc 1799

<210> 2  
<211> 411  
<212> PRT  
<213> human

<220>  
<221> xaa  
<222> 410  
<223> xaa = leu or met

<400> 2  
Met Glu Gln Arg Gly Gln Asn Ala Pro Ala Ala Ser Gly Ala Arg  
1 5 10 15  
Lys Arg His Gly Pro Gly Pro Arg Glu Ala Arg Gly Ala Arg Pro  
20 25 30  
Gly Leu Arg Val Pro Lys Thr Leu Val Leu Val Val Ala Ala Val  
35 40 45  
Leu Leu Leu Val Ser Ala Glu Ser Ala Leu Ile Thr Gln Gln Asp  
50 55 60  
Leu Ala Pro Gln Gln Arg Ala Ala Pro Gln Gln Lys Arg Ser Ser  
65 70 75  
Pro Ser Glu Gly Leu Cys Pro Pro Gly His His Ile Ser Glu Asp  
80 85 90  
Gly Arg Asp Cys Ile Ser Cys Lys Tyr Gly Gln Asp Tyr Ser Thr  
95 100 105  
His Trp Asn Asp Leu Leu Phe Cys Leu Arg Cys Thr Arg Cys Asp  
110 115 120  
Ser Gly Glu Val Glu Leu Ser Pro Cys Thr Thr Thr Arg Asn Thr  
125 130 135  
Val Cys Gln Cys Glu Glu Gly Thr Phe Arg Glu Glu Asp Ser Pro  
140 145 150  
Glu Met Cys Arg Lys Cys Arg Thr Gly Cys Pro Arg Gly Met Val  
155 160 165  
Lys Val Gly Asp Cys Thr Pro Trp Ser Asp Ile Glu Cys Val His  
170 175 180  
Lys Glu Ser Gly Ile Ile Ile Gly Val Thr Val Ala Ala Val Val  
185 190 195  
Leu Ile Val Ala Val Phe Val Cys Lys Ser Leu Leu Trp Lys Lys  
200 205 210  
Val Leu Pro Tyr Leu Lys Gly Ile Cys Ser Gly Gly Gly Asp  
215 220 225

Pro Glu Arg Val Asp Arg Ser Ser Gln Arg Pro Gly Ala Glu Asp  
230 235 240

Asn Val Leu Asn Glu Ile Val Ser Ile Leu Gln Pro Thr Gln Val  
245 250 255

Pro Glu Gln Glu Met Glu Val Gln Glu Pro Ala Glu Pro Thr Gly  
260 265 270

Val Asn Met Leu Ser Pro Gly Glu Ser Glu His Leu Leu Glu Pro  
275 280 285

Ala Glu Ala Glu Arg Ser Gln Arg Arg Arg Leu Leu Val Pro Ala  
290 295 300

Asn Glu Gly Asp Pro Thr Glu Thr Leu Arg Gln Cys Phe Asp Asp  
305 310 315

Phe Ala Asp Leu Val Pro Phe Asp Ser Trp Glu Pro Leu Met Arg  
320 325 330

Lys Leu Gly Leu Met Asp Asn Glu Ile Lys Val Ala Lys Ala Glu  
335 340 345

Ala Ala Gly His Arg Asp Thr Leu Tyr Thr Met Leu Ile Lys Trp  
350 355 360

Val Asn Lys Thr Gly Arg Asp Ala Ser Val His Thr Leu Leu Asp  
365 370 375

Ala Leu Glu Thr Leu Gly Glu Arg Leu Ala Lys Gln Lys Ile Glu  
380 385 390

Asp His Leu Leu Ser Ser Gly Lys Phe Met Tyr Leu Glu Gly Asn  
395 400 405

Ala Asp Ser Ala Xaa Ser  
410